

Claims

1. An isolated nucleic acid molecule selected from the group consisting of
(a) nucleic acid molecules that code for the amino acid sequence of SEQ ID NO:2 or
5 SEQ ID NO:4,
(b) allelic variants of (a), and
(c) complements of (a) or (b).
2. The isolated nucleic acid molecule of claim 1, wherein the isolated nucleic acid
10 molecule codes for SEQ ID NO:2.
3. The isolated nucleic acid molecule of claim 1, wherein the isolated nucleic acid molecule codes for SEQ ID NO:4.
- 15 4. The isolated nucleic acid molecule of claim 1, wherein the isolated nucleic acid molecule comprises the nucleotide sequence of SEQ ID NO:2 or SEQ ID NO:4.
5. An isolated P-glycoprotein polypeptide or fragment thereof which comprises at least one amino acid of a cynomologous P-glycoprotein selected from the group consisting of
20 amino acids 12, 24, 30, 74, 78, 86, 89, 90, 91, 92, 95, 97, 99, 102, 103, 104, 185, 324, 363, 518, 635, 650, 656, 659, 677, 730, 738, 742, 745, 761, 765, 835, 851, 921, 967, 1003, 1027, 1038, 1048, 1103, 1128, 1168 and 1277 of SEQ ID NO:2 and amino acids 93, 94 and 95 of SEQ ID NO:4, wherein the P-glycoprotein is identical to a human P-glycoprotein except for the at least one amino acid of a cynomologous P-glycoprotein
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6. The isolated P-glycoprotein polypeptide or fragment thereof of claim 5, wherein the human P-glycoprotein is selected from the group of SEQ ID NO:5 and SEQ ID NO:6.
7. An isolated P-glycoprotein polypeptide or fragment thereof which comprises at least
30 one amino acid of a cynomologous P-glycoprotein selected from the group consisting of amino acids 3, 6, 8, 10, 13, 17, 19, 20, 21, 26, 30, 36, 38, 48, 52, 56, 64, 74, 78, 84, 85, 86, 87, 88, 89, 90, 91, 92, 93, 94, 95, 96, 98, 100, 101, 102, 103, 104, 105, 106, 110, 113, 145,

190, 197, 210, 231, 319, 324, 327, 345, 363, 395, 451, 455, 456, 468, 473, 494, 518, 530, 631, 641, 642, 648, 650, 655, 656, 664, 665, 672, 673, 674, 675, 683, 687, 689, 691, 692, 694, 701, 705, 715, 729, 730, 734, 742, 743, 745, 754, 757, 765, 835, 912, 918, 921, 940, 941, 944, 966, 967, 968, 970, 972, 981, 1008, 1015, 1023, 1024, 1048, 1093, 1096, 1103, 1128, 1142, 1146, 1147, 1156, 1160, 1163, 1166, 1250 and 1271 of SEQ ID NO:2 and amino acids 93 and 94 of SEQ ID NO:4, wherein the P-glycoprotein is identical to a dog P-glycoprotein except for the at least one amino acid of a cynomolgous P-glycoprotein

8. The isolated P-glycoprotein polypeptide or fragment thereof of claim 7, wherein the dog P-glycoprotein is selected from the group of SEQ ID NO:7 and SEQ ID NO:8.

9. The isolated P-glycoprotein polypeptide or fragment thereof of claim 5 or 7, wherein the amino acid sequence of the polypeptide or fragment thereof is an amino acid sequence selected from the group consisting of SEQ ID NO:2, fragments of SEQ ID NO:2, SEQ ID NO:4 and fragments of SEQ ID NO:4.

10. An isolated nucleic acid molecule which encodes the isolated P-glycoprotein polypeptide or fragment thereof of any of claims 5-9.

11. An expression vector comprising the isolated nucleic acid molecule of claim 1 operably linked to a promoter.

12. An expression vector comprising the isolated nucleic acid molecule of claim 10 operably linked to a promoter.

13. A host cell transformed or transfected with the expression vector of claim 11.

14. A host cell transformed or transfected with the expression vector of claim 12.

15. An agent which selectively binds the isolated polypeptide of claim 5.

16. The method of claim 15, wherein the agent does not bind a human or dog P-

glycoprotein.

17. The agent of claim 15, wherein the agent is a polypeptide.

5 18. The agent of claim 17, wherein the polypeptide is selected from the group consisting of monoclonal antibodies, polyclonal antibodies, Fab antibody fragments, F(ab)₂ antibody fragments and antibody fragments including a CDR3 region.

19. An agent which selectively binds the isolated nucleic acid molecule of claim 1 or
10 claim 10.

20. The agent of claim 19, wherein the agent is an antisense nucleic acid which selectively binds to the isolated nucleic acid molecule.

15 21. A method for predicting the bioavailability of a compound, comprising measuring the transmembrane transport of a test compound by a first P-glycoprotein, comparing the transmembrane transport of the test compound by the first P-glycoprotein and a second P-glycoprotein to predict the bioavailability of the test compound, wherein the relative amount or rate of transport by the first P-glycoprotein and the
20 second P-glycoprotein is predictive of bioavailability of the test compound.

22. The method of claim 21, wherein the first P-glycoprotein is selected from the group consisting of dog P-glycoproteins and primate P-glycoproteins.

25 23. The method of claim 21, wherein the first P-glycoprotein is the polypeptide of claims 5 or 7.

24. The method of claim 21, wherein the second P-glycoprotein is a human P-glycoprotein.

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25. A method for inhibiting P-glycoprotein transporter activity in a mammalian cell comprising

contacting the mammalian cell with an amount of the agent of claim 19 effective to inhibit P-glycoprotein transporter activity in the mammalian cell.

26. A method for increasing bioavailability of a drug in a subject comprising
5 administering to a subject in need of such treatment the agent of claim 19 in an amount effective to increasing bioavailability of a drug.

27. The method of claim 26, wherein the inhibitor is administered prior to administering the drug.

28. The method of claim 26, wherein the inhibitor is administered concurrently with the drug.

29. A method for increasing P-glycoprotein transporter activity in a cell comprising
15 contacting the cell with a molecule selected from the group consisting of the nucleic acid molecule of claim 1 and the nucleic acid molecule of claim 10, in an amount effective to increase P-glycoprotein transporter activity in the cell.

30. A method for identifying lead compounds for a pharmacological agent useful in the
20 treatment of disease associated with P-glycoprotein transporter activity comprising providing a cell or other membrane-encapsulated space comprising a P-glycoprotein as claimed in claim 5 or claim 7;

contacting the cell or other membrane-encapsulated space with a candidate
pharmacological agent under conditions which, in the absence of the candidate
25 pharmacological agent, cause a first amount of P-glycoprotein transporter activity;

determining a second amount of P-glycoprotein transporter activity as a measure of the effect of the pharmacological agent on the P-glycoprotein transporter activity, wherein a second amount of P-glycoprotein transporter activity which is less than the first amount indicates that the candidate pharmacological agent is a lead compound for a pharmacological
30 agent which reduces P-glycoprotein transporter activity and wherein a second amount of P-glycoprotein transporter activity which is greater than the first amount indicates that the candidate pharmacological agent is a lead compound for a pharmacological agent which

increases P-glycoprotein transporter activity.

31. The method of claim 30, further comprising the step of loading the cell or other membrane-encapsulated space with a detectable compound, wherein the compound is detected as a measure of the P-glycoprotein transporter activity.

32. A method for identifying compounds which selectively bind a P-glycoprotein comprising,
contacting the P-glycoprotein claimed in claim 5 or claim 7 with a compound,
determining the binding of the compound to the P-glycoprotein.

33. The method of claim 32 further comprising determining the effect of the compound on the P-glycoprotein transporter activity of the P-glycoprotein.

34. The method of claim 32 further comprising determining the effect of the compound on the ATPase activity of the P-glycoprotein.

35. A method for determining ATPase activity of a P-glycoprotein comprising
contacting the host cell of claim 12 or 14, or a membrane fraction thereof, with
a test drug, and
measuring ATPase activity of the P-glycoprotein.

36. The method of claim 35, wherein the step of measuring ATPase activity is performed at least twice at different times.

37. A method for determining transmembrane transport of a compound by a P-glycoprotein, comprising
contacting the host cell of claim 12 or 14, or a membrane fraction thereof, with
a test drug, and
measuring transport of the test drug under sink conditions in at least one direction of transport selected from the group consisting of the apical to basolateral direction and the basolateral to apical direction.

38. The method of claim 37, wherein the step of measuring transport of the test drug is performed at least twice at different times.